#### STIC-Biotech/ChemLib

From:

Salimi, Ali

Sent:

Thursday, July 18, 2002 2:28 PM STIC-Biotech/ChemLib

To: Subject:

09/514,245

Can you please conduct a search and interference for the application serial no. 09/514,245:

SEQ ID NO: 25

Thanks!

Ali Salimi AU 1648 Room 9D07 Mailbox 8E12

Point of Contact: Susan Hanley
Technical Info. Specialist
CM1 6B05 Tel: 305-4053

Searcher:
Phone:
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Date Picked Up:
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Viruses; ssDNA viruses; Circoviridae;

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Hutet, E. and Albina, E.

Circovirus sequences related to piglet

Patent: WO 9929871-A 12 17-JUN-1999;

HUTET EVELYNE (FR); ALBINA EMMANUEL (F
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Viruses; ssDNA viruses; Circoviridae;

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Hutet, E. and Albina, E.

Circovirus sequences related to pigle!

Patent: WO 9929871-A 9 17-JUN-1999;

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porcine circovirus

Viruses; ssDNA viruses; Circoviridae;

1 (bases 1 to 1767)

Hutet, E. and Albina, E.

Circovirus sequences related to piglet
Patent: WO 9929871-A 10 17-JUN-1999;

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/db_xref="taxon:85708"
/country="France"
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/gene="cap"
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GWLPWDDLLRLCDRYPLTVETKGGTVPFLARSILITSNQTPLEWYSSTAVPAVEALYR
RITSLVFWKNATEQSTEEGGQFVTLSPPCPEFPYEINY"
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Mankertz, A., Domingo, M., Folch, J.M., LeCann, Segales, J., Chmielewicz, B., Plana-Duran, J. a Characterisation of PCV-2 isolates from Spai Virus Res. 66 (1), 65-77 (2000)
20120936
2 (bases 1 to 1767)
Mankertz, A., Domingo, M., Folch, J.M., LeCann, Segales, J., Chmielewicz, B., Plana-Duran, J. a Direct Submission
Submitted (03-NOV-1999) P24, Robert Koch Ins Berlin 13353, Germany
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Pred. No. 2.6e-188;
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/evidence=not_experimental/product="ORF6"
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/note="11.9 kDa"
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Direct Submission

Submitted (04-NOV-1999) Virology and Molecular Epidemiology,

BioScreen European Veterinary Disease Management Center GmbH, 11

Mendelstr., Muenster, NRW 48149, Germany

Location/Qualifiers

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/organism="porcine circovirus type 2"

/db_xref="taxon:85708"
/doutry="Netherlands"

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LLMECGAPRSQGQRSDLSTAVSTLLESGSLVTVAEQHPVTFVRNFRGLAELLKV
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Viruses; ssDNA viruses; Circoviridae; Circovirus.

1 (bases 1 to 1767)

8 Wellenberg, G.J., Pesch, S., Berndsen, F.W., Steverink, P.J.G.M., Hunneman, W., Van der Vorst, T.J.K., Peperkamp, N.H.M.T., Ohlinger, V.F., Schippers, R., Van Oirschot, J.T. and de Jong, M.F. Isolation and characterization of porcine circovirus type 2 fropigs showing signs of post-weaning multisystemic wasting syndro in The Netherlands

Unpublished

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GWLPWDDLLRLCDRYPLTVETKGGTVPFLARSILITSNQTPLEWYSSTAVPAVEALYR
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Meehan, B.M., McNeilly, F., NcNair, I., Walker, I., West, K.,
Ellis, J.A., Krakowka, S., Kennedy, S. and Allan, G.M.
Ellis, J.A., Krakowka, S., Kennedy, S. and Allan, G.M.
Isolation and characterization of porcine circovirus type
Isolation and characterization of porcine dermatitis and nephropat
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1 (bases 1 to 1766)
Meehan, B.M., McNeilly, F., McNair, I., Walker, I., Ellis, Krakowka, S. and Allan, G.M.
Isolation and characterization of porcine circovirus 2 of sow abortion and porcine dermatitis and nephropathy Arch. Virol. 146 (4), 835-842 (2001)
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3 (bases 1 to
Meehan, B.M.
Direct Submission
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Queen's University of Belfast, St
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 Shengbo, C. and Huanchun, Direct Submission Submitted (16-MAY-2001) University, Shi Zi Shan
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/product="replicase"
/protein_id="AAF35300.1
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complement(join(1731. .1767,1. .92))
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                                        complement(1300. .1311)
/note="glycosylation site"
complement(1521. .1610)
/note="ORF6"
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Segales, J., Chmielewicz, B., Plana-Duran, J. and Soil
Characterisation of PCV-2 isolates from Spain, Gery
Virus Res. 66 (1), 65-77 (2000)
2 (bases 1 to 1768)
Mankertz, A., Domingo, M., Folch, J.M., LeCann, P., Jes
Segales, J., Chmielewicz, B., Plana-Duran, J. and Soil
Direct Submission
Submitted (03-NOV-1999) P24, Robert Koch Institut,
Berlin 13353, Germany
Location/Qualifiers
ce /organism="porcine circovirus type 2"
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/db_xref="taxon:85708"
/country="Spain"
complement(37. .738)
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/note="Cap"
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VGEEGNEEGRTPHLQGFANFVKKOTFNKVKWYLGARCHIEKAKGTDQONKEYCSKEGN
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FTPKPVLDSTIDYFQPNNKRNQLWLRLQTSGNVDHVGLGTAFENSKYDQDYNIRVTMY
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                                 Emergence of a new porcine circo Unpublished

2 (bases 1 to 1768)
Wang, L., Willson, P., Chow, B., Gi Direct Submission
Submitted (26-AUG-1998) VIDO, Un Veterinary Road, Saskatoon, Sask Location/Qualifiers

1. 1768
/organism="porcine circ/strain="9741"
/db_xref="taxon:46221"
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/note="putative replication site"
join(1762. .1768,1. .2)
/note="similar to the nonanucleotide circovirus presented in GenBank Acce and U49186"
a 364 c 490 q 465 t
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/codon_start=1
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/COGAPRACOGNETHEOSTEEGGQFVTLSPPCPEFPYEINY"
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Hamel, A.L. and Nayar, G.P.S.
Nucleotide sequence of four d
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Unpublished
3 (bases 1 to 1768)
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Hamel, A.L., Lin, L.L., Sach
PCR detection and characte
Can. J. Vet. Res. 64 (1),
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/rpt_unit=13. .1
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complement(1022. .1027)
complement(1256. .1735)
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YMVYISSSL"
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/codon_start=1
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20120936
2 (bases 1 to 1768)
Mankertz, A., Domingo, M., Folch, J.M., LeCann, P., Jestin, A., Segales, J., Chmielewicz, B., Plana-Duran, J. and Soike, D. Direct Submission
Submitted (03-NOV-1999) P24, Robert Koch Institut, Nordufer 20 Berlin 13353, Germany
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/db_xref="taxon:85708"
/country="Taiwan"
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/organism="porcine
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Lai,S.S.
Direct Submission
Submitted (02-JAN-2002) H
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LLMECGAPRSQGQRSDLSTAVSTLLESGSLVTVAEQHPVTFVRNFRGLAELLKVSGKM
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Imai, K. and Ishihara, R.

Direct Submission

Submitted (27-SEP-2001) Kunitoshi Imai, Hokkaido Research Station,
National Institute of Animal Health, Clinical Microbiology Section,
National Institute of Animal Health, Clinical Microbiology Section,
Hitsujigaoka, Toyohira-ku, Sapporo, Hokkaido 062-0045, Japan
Hitsujigaoka, Toyohira-ku, Sapporo, Hokkaido 062-0045, Japan
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Imai, K., Nishimori, T., Nishimori, K., Ishihara, R. and Sequence of porcine circovirus type 2 from paraffin spigs with postweaning multisystemic wasting sydrome Unpublished
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/rote="viral gene obtained from paraffin-embedded lymph node postweaning multisystemic syndist. 1995
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/specific_host="pig"
/db_xref="taxon:85708"
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#### SUMMARIES

	Description	quence 1, Appl	equence 2, Appl	equence 3, Appl	equence 4, Ap	equence 24, App	equence 2, Appl	equence 38, App	ence 40, App	e 25, App	1, Appl	5, Appl	equence 22, App	equence 2, App	equence 41, App	e 15, A	equence 15, App	equence 7, Appl	e 41, App	equence 13, App	equence 14, App	equence 5, Appl	equence 5, Appl	ce 1, Appl	ď	equence 1, Appl	equence 5, Appl	uence 1, A
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US-09-347-594-1/c

; Sequence 1, Application US/09347594
; Patent No. 6217883
; GENERAL INFORMATION:
; APPLICANT: ALLAN, Gordon M.
; APPLICANT: MEEHAN, Brian M.
; APPLICANT: ELLIS, John A.
; APPLICANT: KRAKOWKA, George S.
; APPLICANT: AUDONNET, Jean-Christophe F.
; TITLE OF INVENTION: PORCINE CIRCOVIRUS AND PARVOVIRUS VACCINE
; FILE REFERENCE: 454313-2338
; CURRENT APPLICATION NUMBER: US/09/347,594
; CURRENT FILING DATE: 1999-07-01
; EARLIER PILING DATE: 1999-07-06
; NUMBER OF SEQ ID NOS: 5
; SOFTWARE: PatentIn Ver. 2.1
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                                                                      Sequence 2, Application US/09347594

Sequence 2, Application US/09347594

Patent No. 6217883

GENERAL INFORMATION:

APPLICANT: ALLAN, Gordon M.

APPLICANT: MEEHAN, Brian M.

APPLICANT: ELLIS, John A.

APPLICANT: ELLIS, John A.

TITLE OF INVENTION: PORCINE CIRCOVIRUS AN FILE OF INVENTION: PORCINE CIRCOVIRUS AN FILE OF INVENTION NUMBER: US/09/347,594

CURRENT FILING DATE: 1999-07-01

EARLIER APPLICATION NUMBER: 98 08777

EARLIER FILING DATE: 1998-07-06

NUMBER OF SEQ ID NOS: 5
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; Sequence 3, Application U
; Patent No. 6217883
; GENERAL INFORMATION:
; APPLICANT: ALLAN, Gordon
; APPLICANT: MEEHAN, Brian
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APPLICANT: ALLAN, Gordon M.
APPLICANT: MEEHAN, Brian M.
APPLICANT: ELLIS, John A.
APPLICANT: ELLIS, John A.
APPLICANT: ELLIS, John A.
APPLICANT: FILIS, John A.
TITLE OF INVENTION: PORCINE CIRCOVIRUS AND PARVOVIRUS VACCINE
FILE REFERENCE: 454313-2338
CURRENT APPLICATION NUMBER: US/09/347,594
CURRENT FILING DATE: 1999-07-01
EARLIER APPLICATION NUMBER: 98 08777
EARLIER FILING DATE: 1998-07-06
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 4
LENGTH: 1768
TYPE: DNA
ORGANISM: Porcine circovirus
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                                               PARVOVIRUS VACCINE
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Pred. No. 4.2e-9;
Mismatches
APPLICANT: ELLIS, John A.
APPLICANT: KRAKOWKA, George S.
APPLICANT: AUDONNET, Jean-Christophe F.
TITLE OF INVENTION: PORCINE CIRCOVIRUS AND PA
FILE REFERENCE: 454313-2338
CURRENT APPLICATION NUMBER: US/09/347,594
CURRENT FILING DATE: 1999-07-01
EARLIER APPLICATION NUMBER: 98 08777
EARLIER FILING DATE: 1998-07-06
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin Ver. 2.1
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Patent No. 6217883
GENERAL INFORMATION:
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Best Local Similarity 92.3%;
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ed. No. 4.5e-86;
Mismatches 222;
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US-09-267-177-38/C
US-09-267-177-38/C
; Sequence 38, Application US/09267177
; Sequence 38, Application US/09267177
; Patent No. 6287856
; GENERAL INFORMATION:
; APPLICANT: Ritchie, Branson W.
; APPLICANT: Ritchie, Branson W.
; APPLICANT: Lukert, Phil D.
; TITLE OF INVENTION: Vaccines against Circovi
; FILE REFERENCE: 21099.0057
; CURRENT APPLICATION NUMBER: 60/077,890
; EARLIER APPLICATION NUMBER: 60/077,890
FARLIER FILING DATE: 1998-03-13
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APPLICANT: Poet, Steven E.
APPLICANT: Ritchie, Branson W.
APPLICANT: Ritchie, Branson W.
APPLICANT: Lukert, Phil D.
TITLE OF INVENTION: Vaccines against Cir
FILE REFERENCE: 21099.0057
CURRENT APPLICATION NUMBER: US/09/267,17; CURRENT FILING DATE: 1999-03-12
EARLIER APPLICATION NUMBER: 60/077,890
EARLIER FILING DATE: 1998-03-13
NUMBER OF SEQ ID NOS: 41
SOFTWARE: FastSEQ for Windows Version 3.
SEQ ID NO 40
LENGTH: 1759
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APPLICANT: Poet, Steven E.

APPLICANT: Ritchie, Branson W.

APPLICANT: Niagro, Frank D.

APPLICANT: Lukert, Phil D.

TITLE OF INVENTION: Vaccines against Cir

FILE REFERENCE: 21099.0057

CURRENT APPLICATION NUMBER: 60/077,890

EARLIER APPLICATION NUMBER: 60/077,890

EARLIER FILING DATE: 1998-03-13

NUMBER OF SEQ ID NOS: 41

SOFTWARE: FastSEQ for Windows Version 3.
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Patent No. 6287856
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ilarity 67.0%;
Conservative
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APPLICANT: Poet, Steven E.
APPLICANT: Ritchie, Branson W.
APPLICANT: Niagro, Frank D.
APPLICANT: Lukert, Phil D.
TITLE OF INVENTION: Vaccines a File Reference: 21099.0057; CURRENT APPLICATION NUMBER: US CURRENT FILING DATE: 1999-03-1; EARLIER APPLICATION NUMBER: 60
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                         48.48; 92.38;
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; Patent No. 6217883
; GENERAL INFORMATION:
; APPLICANT: ALLAN, Gordon M.
; APPLICANT: MEEHAN, Brian M.
; APPLICANT: ELLIS, John A.
; APPLICANT: KRAKOWKA, George S.
; APPLICANT: AUDONNET, Jean-Christophe F.
               Version
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EARLIER FILING DATE: 1998-03-1 NUMBER OF SEQ ID NOS: 41 SOFTWARE: FastSEQ for Windows
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                            LENGTH: 1674
TYPE: DNA
ORGANISM: porcine 0
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IS-09-347-594-5/C
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APPLICANT: Ritchie, Branson W.
APPLICANT: Ritchie, Branson W.
APPLICANT: Lukert, Phil D.
TITLE OF INVENTION: Vaccines against Cirrent APPLICATION NUMBER: US/09/267,17
CURRENT FILING DATE: 1999-03-12
EARLIER APPLICATION NUMBER: 60/077,890
EARLIER FILING DATE: 1998-03-13
NUMBER OF SEQ ID NOS: 41
SOFTWARE: FastSEQ for Windows Version 3.
SEQ ID NO 22
LENGTH: 285
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Patent No. 6287856
GENERAL INFORMATION:
FILE REFERENCE: 454313-2338
CURRENT APPLICATION NUMBER: US/09,
CURRENT FILING DATE: 1999-07-01
EARLIER APPLICATION NUMBER: 98 08
EARLIER FILING DATE: 1998-07-06
NUMBER OF SEQ ID NOS: 5
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 5
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CURRENT APPLICATION NUMBER: US/09/267,17
CURRENT FILING DATE: 1999-03-12
EARLIER APPLICATION NUMBER: 60/077,890
EARLIER FILING DATE: 1998-03-13
NUMBER OF SEQ ID NOS: 41
SOFTWARE: FastSEQ for Windows Version 3
SEQ ID NO 25
LENGTH: 687
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APPLICANT: Poet, Steven E.
APPLICANT: Ritchie, Branson W.
APPLICANT: Niagro, Frank D.
APPLICANT: Lukert, Phil D.
TITLE OF INVENTION: Vaccines against Ciffle Reference: 21099.0057; CURRENT APPLICATION NUMBER: US/09/267,1
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APPLICANT: Poet, Steven E.
APPLICANT: Ritchie, Branson W.
APPLICANT: Ritchie, Branson W.
APPLICANT: Lukert, Phil D.
TITLE OF INVENTION: Vaccines agains: FILE REFERENCE: 21099.0057
CURRENT APPLICATION NUMBER: US/09/24; CURRENT FILING DATE: 1999-03-12; EARLIER APPLICATION NUMBER: 60/077,
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MEDIUM TYPE: Floppy
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California
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LOCATION: 1..322
OTHER INFORMATION:
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           tent No. 6307020
ENERAL INFORMATION:
APPLICANT: Hew, CAPPLICANT: Gong,
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             Score 172.2; DB 4;
Pred. No. 2.8e-44;
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                                                                                                                                                                                                                                  Score 44.6; DB Pred. No. 0.033; ); Mismatches
                                                                                                                                                                                                                                                                                RESULT 13
US-09-103-840A-2/c
; Sequence 2, Application US/09103840A
; Patent No. 6294328
; GENERAL INFORMATION:
; APPLICANT: WHITE, Owen R.
; APPLICANT: WHITE, Owen R.
; APPLICANT: VENTER, Claire M.
; APPLICANT: VENTER, John C.
; TITLE OF INVENTION: DNA SEQUENCES FOR STRAIN
; TITLE OF INVENTION: TUBERCULOSIS
; FILE REFERENCE: 24366-20007.00
; CURRENT APPLICATION NUMBER: US/09/103,840A
; CURRENT FILING DATE: 1998-06-24
; NUMBER OF SEQ ID NOS: 2
; SOFTWARE: PatentIn Ver. 2.1
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ORGANISM: Mycobacterium tuberculosis
FEATURE:
OTHER INFORMATION: CDC 1551
OTHER INFORMATION: "n" bases at var
OTHER INFORMATION: represent a, t, 0
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                    Query Match
Best Local S
Matches 213
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                                                                 l and Crew LL
Eighth Floor
APPLICANT: Gong, Zhiyuan
TITLE OF INVENTION: Intracellular Antifreeze
TITLE OF INVENTION: and Nucleic Acids
NUMBER OF SEQUENCES: 46
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                                                                                                                                                                                                  Version
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Kenneth A.
WIMBER: 31,677
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Mismatches
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30-JAN-1997
INFORMATION:
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                                                                                                                                                 COMPUTER: Floppy disk
COMPUTER: IBM PC COMPATIble
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve
                                                                    Townsend
Center, E
                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/117,12 FILING DATE: 20-NOV-1998 CLASSIFICATION: 435
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TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 41:
SEQUENCE CHARACTERISTICS:
LENGTH: 322 base
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TELECOMMUNICATION INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/246,982A
FILING DATE: May 20, 1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Goldstein, Jorge, A.
REGISTRATION NUMBER: 29,021
REFERENCE/DOCKET NUMBER: 0609.3880002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2540
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 9997 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: linear
                                                                                          Protein And
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Pred. No. 0.0084;
); Mismatches 37;
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                                                                                                                                          Goldstein
                                                                                          DNA,
GENERAL INFORMATION:

APPLICANT: MacDonald, Marcy E.

APPLICANT: Ambrose, Christine M.

APPLICANT: Duyao, Mabel P.

APPLICANT: Gusella, James F.

TITLE OF INVENTION: Huntingtin DNZ
NUMBER OF SEQUENCES: 25

CORRESPONDENCE ADDRESS:

ADDRESSEE: Sterne, Kessler, Golc
STREET: 1100 New York Avenue
CITY: Washington
STATE: D.C.

COUNTRY: U.S.A.

ZIP: 20005
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US-08-246-982A-15
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Compugen Ltd
version - 2000
 GenCore
(c) 1993
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nucleic search, using sw model OM nucleic

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atgacgtatccaaggaggcg Title: Perfect score: Sequence:

Gapext IDENTITY\_NUC Gapop 10.0 , Scoring table:

1.0

residues 13736207 seqs, 6748477542 Searched:

hits satisfying chosen parameters: of number Total

seq length: 0 seq length: 2000000000 Minimum DB Maximum DB

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

EST:\*

em\_estba:\*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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BASE COUNT ORIGIN

293059 GH16139. G445569 GAEa0 402428 GH21648. 530753 vj44e07. 296628 LP10591. I619435 RH50469	1019848 KH309/1. 857154 603007B08 K020691 Mus musc 589477 vl47h12.s 798859 vv94a07.r	BG591496 EST499338 AA698799 HL05714.5 BG599848 EST504743 BM110666 EST558202 BF253141 EST445636 AL151849 Anopheles AW615354 hh74b04.x	I141028 oy68h12. I364938 qz23d02. BI799844 H142B04 C99980 C99980 Or I017603 ou29e09. BI561078 6032542	181814 AU18181 238831 GH14954 1614132 RH4343 1750241 FG02_0
2 554 9 AI29305 2 870 10 BG4455 2 479 9 AI40242 2 521 9 AA53075 2 555 9 AI29662 2 574 10 BI6194	2 591 9 AI857154 2 835 11 AK02069 1 452 9 AA589477 1 514 9 AA798859	1 617 10 1 636 9 1 650 10 1 706 10 1 710 10 1 1064 12	1 494 9 AI14102 1 496 9 AI36493 1 574 10 BI7998 1 575 10 C99980 1 592 9 AI01760 1 657 10 BI5610	459 9 AU1818 580 9 AI2388 653 10 BI614 655 10 BI750
44 44 43 .8 8 6 6 9 8 4 4 4 8 .2 8 8 6 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9	43.2 43.2 43.2 6 43.0 6 6 6	29 43 6. 30 43 6. 31 43 6. 32 43 6. 34 42.8 6. 35 42.6 6.	6 42.6 6 8 42.6 6 9 42.6 6 0 42.6 6	2 4 4 4 4 2 4 2 4 2 4 2 4 2 4
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## ALIGNMENTS

A1514058 GH27112.5prime GH Drosophila melanogaster head por2 Drosophila melanogaster cDNA clone GH27112 5prime, mRNA sequence. A1514058 A1514058 A1514058.1 GI:4418120 EST. fruit fly. Drosophila melanogaster Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Ephydroidea; Drosophilac. 1 (bases I to 561) Harvey,D., Brokstein,P., Hong,L., Evans-Holm,M., Su,C., Tsang,G., Lewis,S. and Rubin,G.M. BDGP/HHMI Drosophila EST Project Unpublished (2001) Contact: Stapleton, M. BDGP Lawrence Berkeley National Lab One Cyclotron Rd, Berkeley, CA 94720, USA Fax: 510 486 6798 Email: http://www.fruitfly.org/EST, est@fruitfly.berkeley.edu Plate: 271 row: A column: 12 High quality sequence stop: 488. Location/Qualifiers	1561 /organism="Drosophila melanogaster" /db_xref="taxon:7227" /clone="GH27112" /clone_lib="GH Drosophila melanogaster head pOT2" /sex="male and female" /dev_stage="adult" /lab_host="DH5 - alpha" /note="Organ: head; Vector: pOT2; Site_1: EcoRI; Site_2: xhoI; Sized fractionated cDNAs were directly ligated into pOT2. Plasmid cDNA library."	$\simeq$
RESULT 1 A1514058/C LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM TITLE JOURNAL COMMENT FEATURES	source	BASE COUNT

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ORIGIN
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S Genoscope.

Direct Submission

L Submitted (02-JUN-1999) Genoscope - Centre National de Sequencage:
Direct Submission

L Submitted (02-JUN-1999) Genoscope - Centre National de Sequencage:
BP 191 91006 EVRY cedex - FRANCE (E-mail: segref@genoscope.cns.fr

- Web: www.genoscope.cns.fr)

Determination of this BAC-end sequence was carried out as part of a collaboration with the Berkeley Drosophila Genome Project (BDGP).
The BDGP is constructing a physical map of the Drosophila melanogaster genome using these BACs. For further information please see http://www.fruitfly.org The BDGP Drosophila melanogaster BAC library was prepared by Kazutoyo Osoegawa and Aaron Mammoser in Pieter de Jong's laboratory in the Department of Cancer Genetics at the Roswell Park Cancer Institute in Buffalo, NY. The library is named RPCI-98 and was constructed by partial ECORI digestion of Drosophila DNA provided by the BDGP from the isogenic strain y2; cn bw sp, the same strain used for the library and how to order individual BAC clones, the entire library, or filters for hybridization from the BACPAC Resource Center can be found at http://bacpac.med.buffalo.edu/drosophila_bac.htm.

Location/Qualifiers

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Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Ins
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
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/db_xref="taxon:7227"
/clone_lib="RPCI-98"
/clone="BACR26H16"
/note="end: TET3"
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Mismatches
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Pred. No. 1
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larity 18.1%;
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Similarity 58.5%;
9; Conservative
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AUTHORS
TITLE
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B1629442
RH58310.5prime RH Drosophila melanogaster normalized Head pFlc-1
Drosophila melanogaster cDNA clone RH58310 5 similar to CG17108:
FBan0017108 GO:[] located on: 2L 32Al-32Al;: 08/23/2001, mRNA
sequence.
B1629442
B1629442.1 GI:15531652
EST.
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Stapleton, M., Brokstein, P., Hong, L., Tyler, D., Berman, B., Carlson, J., Champe, M., Chavez, C., Dorsett, V., Farfan, D., Frise, E., Georg, R., Gonzalez, M., Guarin, H., Harris, N., Li, P., Liao, G., Misra, S., Mungall, C.J., Nunoo, J., Pacleb, J., Paragas, V., Park, S., Phouanenavong, S., Wan, K., Yu, C., Lewis, S.E., Celniker, S. and Rubi
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127 c 269 g 153 t 2 others
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One Cyclotron Rd, Berkeley, CA 94720, USA
Fax: 510 486 6798
Email: http://www.fruitfly.org/EST, est@fruitfly.berkeley.edu
hit genomic AE003629: arm:2L [10413221,10674011]
estimated-cyto:31E5-32A5: 08/23/2001
Plate: RH.583 row: A column: 10
High quality sequence stop: 511.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        fruit fly.
Drosophila melanogaster
Drosophila melanogaster
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Eukaryota; Metazoa; Arthropoda; Tracheata; Brachycera;
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.

Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.

1 (bases 1 to 638)

1 (bases 1 to 638)
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BDGP/HHMI RH Drosophila EST Project
Unpublished (2001)
Contact: Stapleton, M.
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72 a 91 c 224 g 106 t
                                                 Tsang, G
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GH14332.5prime GH Drosophila melanogaster head pOT2 Drosophila melanogaster cDNA clone GH14332 5prime, mRNA sequence.
A1238309.1 GI:3833167
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Drosophila melanogaster
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Ins
Pterygota; Metazoa; Arthropoda; Tracheata; Brachycera;
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
1 (bases 1 to 497)
Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C.,
Lewis, S. and Rubin, G.M.
BDGP/HHMI Drosophila EST Project
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          era; Brachyc
Drosophila.
Pterygota; Neophere,
Muscomorpha; Ephydroidea; Drosephila,
1 (bases 1 to 493)
Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su Lewis, S. and Rubin, G.M.
BDGP/HHMI Drosophila EST Project
Unpublished (2001)
Contact: Stapleton, M.
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Pred. No. 1.9;
); Mismatches
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                                                                                                                           Lawrence Berkeley National Lab
One Cyclotron Rd, Berkeley, CA 94720,
Fax: 510 486 6798
Email: http://www.fruitfly.org/EST, e
Plate: 258 row: G column: 7
High quality sequence stop: 492.
Location/Qualifiers
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One Cyclotron Rd, Berkeley, CA 94720
Fax: 510 486 6798
Email: http://www.fruitfly.org/EST,
Plate: 143 row: C column: 8
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ilarity 58.3%;
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pOT2. Plasmid cDNA library."
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GH12051.5prime GH Drosophila melanogaster head pOT2 Drosophila melanogaster cDNA clone GH12051 5prime, mRNA sequence.
AI134557.1 GI:3627115
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Drosophila melanogaster

Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insections of the second of the second
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One Cyclotron Rd, Berkeley, CA 94720,
Fax: 510 486 6798
Email: http://www.fruitfly.org/EST, es
Plate: 120 row: E column: 3
High quality sequence stop: 403.
Location/Qualifiers
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Similarity 58.3%;
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DB 10;

EST 19-APR Drosophila

mRNA linear E3 anogaster head pOT2 D3 5prime, mRNA sequence

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est@fruitfly.berkeley.edu

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xhoI; Sized fractionated cDNAs were directly ligated into pOT2. Plasmid cDNA library."

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                                                                                                                                                                                               Drosophila melanogaster

Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insect
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.

E 1 (bases 1 to 484)
5 Harvey,D., Brokstein,P., Hong,L., Evans-Holm,M., Su,C., Tsa
Lewis,S. and Rubin,G.M.
BDGP/HHMI Drosophila EST Project
Unpublished (2001)
Contact: Stapleton, M.
BDGP
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One Cyclotron Rd, Berkeley, CA 94720,
Fax: 510 486 6798
Email: http://www.fruitfly.org/EST, e:
Plate: 70 row: E column: 10
High quality sequence stop: 404.
Location/Qualifiers
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Pred. No. 1.9;
0; Mismatches
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GH07058.5prime GH Drosophila melanogaster cDNA clone GH07058 5p AI108264
AI108264.1 GI:3476543
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ilarity 58.3%;
Conservative
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GH25879.5prime (melanogaster cD)
A1405757
A1405757.1 GI:
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Best Local S
Matches 77
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                                                              RESULT 5
AI108264/c
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/db_xref="taxon:7227"
/clone="GH15617"
/clone_lib="GH Drosophila melanogaster head pOT2"
/sex="male and female"
/sex="male and female"
/dev_stage="adult"
/lab_host="DH5 - alpha"
/lab_host="DH5 - alpha"
/note="Organ: head; Vector: pOT2; Site_1: EcoRI; Site_2: xhoI; Sized fractionated cDNAs were directly ligated intoportational potautional po
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Lawrence Berkeley National Lab
One Cyclotron Rd, Berkeley, CA 94720, USA
Fax: 510 486 6798
Email: http://www.fruitfly.org/EST, est@fruitfly.berkeley.edu
Plate: 156 row: B column: 5
High quality sequence stop: 293.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           fruit fly.

Drosophila melanogaster
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.

E 1 (bases 1 to 445)
S Harvey,D., Brokstein,P., Hong,L., Evans-Holm,M., Su,C., Tsan Lewis,S. and Rubin,G.M.
BDGP/HHMI Drosophila EST Project
Unpublished (2001)
Contact: Stapleton, M.
BDGP
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                                       59;
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Pred. No. 1.9;
0; Mismatches
Score 44.2; D)
Pred. No. 1.8;
                         d. No. 1.8
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larity 57.2%;
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Best Local Similarity 58.3%;
Matches 77; Conservative
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GH15617.5prime GH E
melanogaster cDNA c
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AI292664.1 GI:3942
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                    . Similarity 79; Conser
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Best Local S
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/lab_host="DH5 - alpha"
/note="Organ: head; Vector: pOT2; Site_1: EcoRI; Site_
XhoI; Sized fractionated cDNAs were directly ligated in
pOT2. Plasmid cDNA library."
109 c 241 g 133 t
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                                                                                  Tsang,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Insecta;
                                                                                                                                                                                                               est@fruitfly.berkeley.edu
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1 (bases 1 to 561)
Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C., Tewis, S. and Rubin, G.M.

BDGP/HHMI Drosophila EST Project
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0
          melanogaster
Metazoa; Arthropoda; Tracheata; Hexapoda; In:
Neoptera; Endopterygota; Diptera; Brachycera
a: Ephydroidea; Drosophilidae; Drosophila.
                                                                                   Su, C.,
                                                                                                                                                                                                                                                                                                                               head
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                                                                                   Evans-Holm, M.,
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One Cyclotron Rd, Berkeley, CA 94720, USA
                                                                                                                                                                                    USA
                                                                                                                                                                  Lawrence Berkeley National Lab
One Cyclotron Rd, Berkeley, CA 94720,
Fax: 510 486 6798
Email: http://www.fruitfly.org/EST, e:
Plate: 146 row: C column: 8
High quality sequence stop: 489.
Location/Qualifiers
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                                                                                                                                                                                   94720,
                                                                                                                                                                                                                                                                    1. .558
/organtsm="Drosophtla melan/db_xref="taxon:7227"
/clone="GH14632"
/clone_lib="GH Drosophila m/sex="male and female"
/dev_stage="adult"
/acv_stage="adult"
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                                               Muscomorpha; Ephydroidea; Drosophil (bases 1 to 558)
Harvey, D., Brokstein, P., Hong, L., Lewis, S. and Rubin, G.M.
BDGP/HHMI Drosophila EST Project Unpublished (2001)
Contact: Stapleton, M.
BDGP
                                                                                   Hong, L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 44;
Pred. No.
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Drosophila melanogaster
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illarity 58.3%;
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Local Similarity
hes 77; Conserv
fruit fly.
Drosophila
Eukaryota;
Pterygota;
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AUTHORS
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561 bp mRNA linear EST 19-APR-2001 Drosophila melanogaster head pOT2 Drosophila clone GH26478 5prime, mRNA sequence.
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/lab_host="DH5 - alpha"
/note="Organ: head; Vector: pOT2; Site_1: EcoRI; Site_2:
XhoI; Sized fractionated cDNAs were directly ligated into pOT2. Plasmid cDNA library."
110 c 245 g 132 t
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1 (bases 1 to 561)
Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C., Tsanglewis, S. and Rubin, G.M.
BDGP/HHMI Drosophila EST Project
Unpublished (2001)
Contact: Stapleton, M.
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                                                                                                                  /organism="Drosophila melanogaster"
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/clone="GH24758"
/clone_lib="GH Drosophila melanogaster
/sex="male and female"
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/clone="GH26478"
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One Cyclotron Rd, Berkeley, CA 94720,
Fax: 510 486 6798
Email: http://www.fruitfly.org/EST, e
Plate: 264 row: G column: 6
High quality sequence stop: 494.
Location/Qualifiers
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Fax: 510 486 6798
Email: http://www.fruitfly.org/EST,
Plate: 247 row: E column: 10
High quality sequence stop: 461.
Location/Qualifiers
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/sex="male and female"
/dev_stage="adult"
/lab_host="DH5 - alpha"
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/db_xref="taxon:7227"
/clone="GH23705"
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/note="Organ: head; Vector: pOT2; Site_1: EcoRI; Site_2: xhoI; Sized fractionated cDNAs were directly ligated int pOT2. Plasmid cDNA library."
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Drosophila melanogaster

Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;

Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;

Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.

1 (bases 1 to 557)

Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C., Tsang Lewis, S. and Rubin, G.M.

BDGP/HHMI Drosophila EST Project
Unpublished (2001)

Contact: Stapleton, M.
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GH Drosophila melanogaster head pOT2 D1
NNA clone GH23705 5prime, mRNA sequence
                                Length 532
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One Cyclotron Rd, Berkeley, CA 94720,
Fax: 510 486 6798
Email: http://www.fruitfly.org/EST, er
Plate: 237 row: A column: 5
High quality sequence stop: 455.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                94720,
                               Score 44; DB 9
Pred. No. 1.9;
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Pterygota; |
                                          Similarity 77; Conser
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557 bp mRNA linear EST 19-APR-2001 e GH Drosophila melanogaster head pOT2 Drosophila cDNA clone GH26313 5prime, mRNA sequence.
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Drosophila
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/lab_host="DH5 - alpha"
/note="Organ: head; Vector: pOT2; Site_1: EcoRI; Site_2:
XhoI; Sized fractionated cDNAs were directly ligated into pOT2. Plasmid cDNA library."
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Drosophila melanogaster
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Ins
Pterygota; Metazoa; Arthropoda; Tracheata; Brachycera;
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.

1 (bases 1 to 557)
Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C.,
Lewis, S. and Rubin, G.M.
BDGP/HHMI Drosophila EST Project
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Drosophila melanogaster head pOT2 D1
clone GH14632 5prime, mRNA sequence
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One Cyclotron Rd, Berkeley, CA 94720,
Fax: 510 486 6798
Email: http://www.fruitfly.org/EST, ex
Plate: 263 row: B column: 1
High quality sequence stop: 494.
Location/Qualifiers
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/organism="Drosophila meland/
/db_xref="taxon:7227"
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/clone_lib="GH Drosophila m
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ilarity 58.3%;
Conservative
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Contact: Stapleton,
BDGP
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Job time: 6609 sec
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E 1 (bases 1 to 575)
S Harvey, D., Brokstein, P., Hong, L., Evans-Holm, M., Su, C., Tsang, G. Lewis, S. and Rubin, G.M.
BDGP/HHMI Drosophila EST Project
Unpublished (2001)
Contact: Stapleton, M.
BDGP
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/note="Organ: head; Vector: pOT2; Site_1: EcoRI; Site_3
/note="Organ: head; Vector: pOT2; Site_1: EcoRI; Site_3
XhoI; Sized fractionated cDNAs were directly ligated in pOT2; Plasmid cDNA library."
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/db_xref="taxon:7227"
/clone="GH22602"
/clone_lib="GH Drosophila melanogaster
/sex="male and female"
XhoI; Sized fractionated cDNAs were pOT2. Plasmid cDNA library."
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One Cyclotron Rd, Berkeley, CA 94720,
Fax: 510 486 6798
Email: http://www.fruitfly.org/EST, eplate: 226 row: A column: 2
High quality sequence stop: 437.
Location/Qualifiers
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Pterygota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophiliae; Drosophila.
E 1 (bases 1 to 593)
S Harvey,D., Brokstein,P., Hong,L., Evans-Holm,M., Su,C., Tsang,G. Lewis,S. and Rubin,G.M.
BDGP/HHMI Drosophila EST Project
Unpublished (2001)
Contact: Stapleton, M.
BDGP
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/organism="Drosophila melanogaster"
/db_xref="taxon:7227"
/clone="GH27392"
/clone_lib="GH Drosophila melanogaster he
/sex="male and female"
/dev_stage="adult"
/lab_host="DH5 - alpha"
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One Cyclotron Rd, Berkeley, CA 94720,
Fax: 510 486 6798
Email: http://www.fruitfly.org/EST, e:
Plate: 273 row: H column: 8
High quality sequence stop: 547.
Location/Qualifiers
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residues 858457221 seds, 1736436 Searched

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Database

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#### SUMMARIES

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Hassard L;
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Haines D, Krakowka GS,

Allan GM, Meehan B, Clark E, Charreyre CE, Chappuis GE,

Ellis JA, A Harding J, Mcneilly F;

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### ALIGNMENTS

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ALVAC; porcine circovirus-2; vaccine; pig; myocarditis; abortion; intrauterine infection; multisystemic wasting syndrome; ds.
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                  standard; DNA;
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                                                         The present invention relates to the use of porcine circovirus-2 (PCV-; immunogen to formulate a vaccine composition to prevent or treat pigs against myocarditis and/or abortion and/or intrauterine infection and/o post-weaning, multisystemic wasting syndrome and other pathological sequalae associated with PCV-2. The present sequence is a DNA fragment a plasmid, which expresses gene products of PCV-2.
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                                                                                                                                                                                                                                              el recombinant virus comprising DNA from F
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The present invention describes a new isolated porcine circovirus Type II (PCVII), obtained from postweaning multisystemic wasting syndrome-affected pigs. AAX83754 to AAX83757 represent PCVII nucleotide sequences. AAY24929 to AAY24934 represent PCVII open reading frame (ORF) proteins (N.B. the PCVII ORFs given in Fig 2A to Fig 2B do not correspond exactly with the PCVII ORFs given in Fig 3A to Fig 3D). The PCVII polypeptides can be used for treating or preventing PCVII infection in vertebrates. The products can also be used to detect the
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The present invention relates to the use of porcine circovirus-2 (PCV-2) immunogen to formulate a vaccine composition to prevent or treat pigs against myocarditis and/or abortion and/or intrauterine infection and/or post-weaning, multisystemic wasting syndrome and other pathological sequalae associated with PCV-2. The present sequence is a DNA fragment or a strain of PCV, which was used in the present invention.
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sequence is a DNA fragment
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The present invention describes a new isolated porcine circovirus Type II (PCVII), obtained from postweaning multisystemic wasting syndrome-affected pigs. AAX83754 to AAX83757 represent PCVII nucleotide equences. AAY24929 to AAY24934 represent PCVII open reading frame (ORF) roteins (N.B. the PCVII ORFs given in Fig 2A to Fig 2B do not
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The present invention describes a new isolated porcine circovirus Type II (PCVII), obtained from postweaning multisystemic wasting syndrome-affected pigs. AAX83754 to AAX83757 represent PCVII nucleotide sequences. AAY24929 to AAY24934 represent PCVII open reading frame (ORI proteins (N.B. the PCVII ORFs given in Fig 2A to Fig 2B do not correspond exactly with the PCVII ORFs given in Fig 3A to Fig 3D). The PCVII polypeptides can be used for treating or preventing PCVII infection in vertebrates. The products can also be used to detect the PCVII.
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The present sequence represents the nucleotide sequence of PCV isolate Impl011-48121. The specification describes a preparation of type II porcine circovirus (PCV), which is particularly isolated from a lesion, from a pig with symptoms of PMWS (porcine multisystemic wasting syndrome). PCV (attenuated or inactivated), polypeptides derived from it, and vectors that express these polypeptides are all useful in vaccines, suitable for administration to adult or young pigs, or to pregnant sows (for passive immunization of their offspring). DNA isolated from PCV is used for in vivo or in vitro expression of viral polypeptides, also as probes or primers for diagnosis in usual hybridization or amplification assays. These polypeptides may also be used diagnostically to detect PCV-specific antibodies, while antibodies raised against the polypeptides can be used to detect antigens, in any
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The invention provides a novel antigenic preparation directed against porcine multisystemic wasting syndrome (PMWS) that comprises porcine circovirus (PCV) antigen and porcine parvovirus (PPV) antigen. The PCV antigens can be used for vaccination against PMWS. The present sequence represents the DNA sequence of PCV Imp.1011-48121 isolate.
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                                                                                                                       The present invention relates to the use of porcine circovirus-2 (PCV-immunogen to formulate a vaccine composition to prevent or treat pigs against myocarditis and/or abortion and/or intrauterine infection and/o post-weaning, multisystemic wasting syndrome and other pathological sequalae associated with PCV-2. The present sequence is a DNA fragment a strain of PCV, which was used in the present invention.
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Pred. No. 4.2e-105;
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The present sequence represents the nucleotide sequence of PCV isolate Imp1011-48285. The specification describes a preparation of type II porcine circovirus (PCV), which is particularly isolated from a lesion, from a pig with symptoms of PMWS (porcine multisystemic wasting syndrome). PCV (attenuated or inactivated), polypeptides derived from it, and vectors that express these polypeptides are all useful in vaccines, suitable for administration to adult or young pigs, or to pregnant sows (for passive immunization of their offspring). DNA isolated from PCV is used for in vivo or in vitro expression of viral polypeptides, also as probes or primers for diagnosis in usual hybridization or amplification assays. These polypeptides may also be used diagnostically to detect PCV-specific antibodies, while antibodies raised against the polypeptides can be used to detect antigens, in any
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Pred. No. 3.9e
0; Mismatches
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No. 3.
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ilarity 98.5%;
Conservative
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97FR-0012382.
98FR-0000873.
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03-OCT-1997;
22-JAN-1998;
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The present sequence represents the nucleotide sequence of PCV isolate Imp1011-48285. The specification describes a preparation of type II porcine circovirus (PCV), which is particularly isolated from a lesion, from a pig with symptoms of PMWS (porcine multisystemic wasting syndrome). PCV (attenuated or inactivated), polypeptides derived from 1t, and vectors that express these polypeptides are all useful in vaccines, suitable for administration to adult or young pigs, or to pregnant sows (for passive immunization of their offspring). DNA isolated from PCV is used for in vivo or in vitro expression of viral polypeptides, also as probes or primers for diagnosis in usual hybridization or amplification assays. These polypeptides may also be used diagnostically to detect PCV-specific antibodies, while antibodies raised against the polypeptides can be used to detect antigens, in any usual immunoassay format.
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